

M Q R L G A T L L C																			10
GGCACGAGGGGGCGGCGCTGCGGGGCGCAGAGCGGAG ATG CAG CGG CTT GGG GCC ACC CTG CTG TGC																			67
L L L A A A V P T A P A P A P T A T S A	30																		
CTG CTG CTG GCG GCG GCG GTC CCC ACG GCC CCC GCG CCC GCT CCG ACG GCG ACC TCG GCT	127																		
P V K P G P A L S Y P Q E E A T L N E M	50																		
CCA GTC AAG CCC GGC CCG GCT CTC AGC TAC CCG CAG GAG GAG GCC ACC CTC AAT GAG ATG	187																		
F R E V E E L M E D T Q H K L R S A V E	70																		
TTC CGC GAG GTT GAG GAA CTG ATG GAG GAC ACG CAG CAC AAA TTG CGC AGC GCG GTG GAA	247																		
E M E A E E A A A K A S S E V N L A N L	90																		
GAG ATG GAG GCA GAA GAA GCT GCT GCT AAA GCA TCA TCA GAA GTG AAC CTG GCA AAC TTA	307																		
P P S Y H N E T N T D T N V G N N T I H	110																		
CCT CCC AGC TAT CAC AAT GAG ACC AAC ACA GAC ACG AAC GTT GGA AAT AAT ACC ATC CAT	367																		
V H R E I H K I T N N Q T G Q M V F S E	130																		
GTG CAC CGA GAA ATT CAC AAG ATA ACC AAC AAC CAG ACT GGA CAA ATG GTC TTT TCA GAG	427																		
T V I T S V G D E E G R R S H E C I I D	150																		
ACA GTT ATC ACA TCT GTG GGA GAC GAA GAA GGC AGA AGG AGC CAC GAG TGC ATC ATC GAC	487																		
E D C G P S M Y C Q F A S F Q Y T C Q P	170																		
GAG GAC TGT GGG CCC AGC ATG TAC TGC CAG TTT GCC AGC TTC CAG TAC ACC TGC CAG CCA	547																		
C R G Q R M L C T R D S E C C G D Q L C	190																		
TGC CGG GGC CAG AGG ATG CTC TGC ACC CGG GAC AGT GAG TGC TGT GGA GAC CAG CTG TGT	607																		
V W G H C T K M A T R G S N G T I C D N	210																		
GTC TGG GGT CAC TGC ACC AAA ATG GCC ACC AGG GGC AGC AAT GGG ACC ATC TGT GAC AAC	667																		
Q R D C Q P G L C C A F Q R G L L F P V	230																		
CAG AGG GAC TGC CAG CCG GGG CTG TGC TGT GCC TTC CAG AGA GGC CTG CTG TTC CCT GTG	727																		
C T P L P V E G E L C H D P A S R L L D	250																		
TGC ACA CCC CTG CCC GTG GAG GGC GAG CTT TGC CAT GAC CCC GCC AGC CGG CTT CTG GAC	787																		
L I T W E L E P D G A L D R C P C A S G	270																		
CTC ATC ACC TGG GAG CTA GAG CCT GAT GGA GCC TTG GAC CGA TGC CCT TGT GCC AGT GGC	847																		
L L C Q P H S H S L V Y V C K P T F V G	290																		
CTC CTC TGC CAG CCC CAC AGC CAC AGC CTG GTG TAT GTG TGC AAG CCG ACC TTC GTG GGG	907																		
S R D Q D G E I L L P R E V P D E Y E V	310																		
AGC CGT GAC CAA GAT GGG GAG ATC CTG CTG CCC AGA GAG GTC CCC GAT GAG TAT GAA GTT	967																		
G S F M E E V R Q E L E D L E R S L T E	330																		
GGC AGC TTC ATG GAG GAG GTG CGC CAG GAG CTG GAG GAC CTG GAG AGG AGC CTG ACT GAA	1027																		
E M A L L E P A A A A A A L L G R E E I	350																		
GAG ATG GCG CTG AGG GAG CCT GCG GCT GCC GCC GCT GCA CTG CTG GGA AGG GAA GAG ATT	1087																		
TAG																			351 1090

Figure 1B

ATCTGGACCAGGCTGTGGGTAGATGTGCAATAGAAATAGCTAATTTATTTCCCCANGTGTGTGCTTTAAGCGTGGGCTG 1169
ACCAGGCTTCTTCCTACATCTTCTTCCAGTAAGTTTCCCCTCTGGCTTGACAGCATGAGGTGTTGTGCATTTGTTTCAG 1248
CTCCCCCAGGCTGTTCTCCAGGCTTCACAGTCTGGTGTCTGGGAGAGTCAGGCAGGGTTAAACTGCAGGAGCAGTTTGC 1327
CACCCCTGTCCAGATTATTGGCTGTCTTGCCTCTACCAGTTGGCAGACAGCCGTTTGTCTACATGGCTTTGATAATTG 1406
TTTGAGGGGAGGAGATGGAAACAATGTGGAGTCTCCCTCTGATTGGTTTTTGGGGAAATGTGGAGAAGAGTGCCCTGCTT 1485
TGCAAACATCAACCTGGCAAAAATGCAACAAATGAATTTTCCACGCAGTTCTTTCCATGGGCATAGGTAAGCTGTGCCT 1564
TCAGCTGTTGCAGATGAAATGTTCTGTTCACCCTGCATTACATGTGTTTATTTCATCCAGCAGTGTGCTCAGCTCCTAC 1643
CTCTGTGCCAGGGCAGCATTTTCATATCCAAGATCAATTCCTCTCTCAGCACAGCCTGGGGAGGGGGTCATTGTTCTC 1722
CTCGTCCATCAGGGATTTTCAGAGGCTCAGAGACTGCAAGCTGCTTGCCCAAGTCACACAGCTAGTGAAGACCAGAGCAG 1801
TTTCATCTGGTTGTGACTCTAAGCTCAGTGTCTCTCTCACTACCCACACCAGCCTTGGTGCCACCAAAGTGCTCCCC 1880
AAAAGGAAGGAGAATGGGATTTTCTTTTGAGGCATGCACATCTGGAATTAAGGTCAAACCTAATTCTCACATCCCTCTA 1959
AAAGTAACTACTGTTAGGAACAGCAGTGTCTCACAGTGTGGGGCAGCCGTCCTTCTAATGAAGACAATGATATTGAC 2038
ACTGTCCCTCTTTGGCAGTTGCATTAGTAACTTTGAAAGGTATATGACTGAGCGTAGCATACAGGTTAACCTGCAGAAA 2117
CAGTACTTAGGTAATTGTAGGGCGAGGATTATAAATGAAATTTGCAAAATCACTTAGCAGCAACTGAAGACAATTATCA 2196
ACCACGTGGAGAAAATCAAACCGAGCAGGGCTGTGTGAAACATGGTTGTAATATGCGACTGCGAACACTGAACTCTACG 2275
CCACTCCACAAATGATGTTTTTCAGGTGTCATGGACTGTTGCCACCATGTATTTCATCCAGAGTTCTTAAAGTTTAAAGTT 2354
GCACATGATTGTATAAGCATGCTTTCTTTGAGTTTTAAATTATGTATAAACATAAGTTGCATTTAGAAATCAAGCATAA 2433
ATCACTTCACTGCTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 2479

Figure 2

GAATTCGGCACGAGAGACGACGTGCTGAGCTGCCAGCTTAGTGGAAGCTCTGCTCTGGGTGGAGAGCAGCCTCGCTTTG	79
GTGACGCACAGTGCTGGGACCCTCCAGGAGCCCCGGGATTGAAGG	8
L S W L C S P L G A L V L D F N N I R S	148
CTG AGC TGG CTC TGC TCT CCC CTG GGA GCT CTG GTC CTG GAC TTC AAC AAC ATC AGG AGC	28
S A D L H G A R K G S Q C L S D T D C N	208
TCT GCT GAC CTG CAT GGG GCC CGG AAG GGC TCA CAG TGC CTG TCT GAC ACG GAC TGC AAT	48
T R K F C L Q P R D E K P F C A T C R G	268
ACC AGA AAG TTC TGC CTC CAG CCC CGC GAT GAG AAG CCG TTC TGT GCT ACA TGT CGT GGG	68
L R R R C Q R D A M C C P G T L C V N D	328
TTG CGG AGG AGG TGC CAG CGA GAT GCC ATG TGC TGC CCT GGG ACA CTC TGT GTG AAC GAT	88
V C T T M E D A T P I L E R Q L D E Q D	388
GTT TGT ACT ACG ATG GAA GAT GCA ACC CCA ATA TTA GAA AGG CAG CTT GAT GAG CAA GAT	108
G T H A E G T T G H P V Q E N Q P K R K	448
GGC ACA CAT GCA GAA GGA ACA ACT GGG CAC CCA GTC CAG GAA AAC CAA CCC AAA AGG AAG	128
P S I K K S Q G R K G Q E G E S C L R T	508
CCA AGT ATT AAG AAA TCA CAA GGC AGG AAG GGA CAA GAG GGA GAA AGT TGT CTG AGA ACT	148
F D C G P G L C C A R H F W T K I C K P	568
TTT GAC TGT GGC CCT GGA CTT TGC TGT GCT CGT CAT TTT TGG ACG AAA ATT TGT AAG CCA	168
V L L E G Q V C S R R G H K D T A Q A P	628
GTC CTT TTG GAG GGA CAG GTC TGC TCC AGA AGA GGG CAT AAA GAC ACT GCT CAA GCT CCA	188
E I F Q R C D C G P G L L C R S Q L T S	688
GAA ATC TTC CAG CGT TGC GAC TGT GGC CCT GGA CTA CTG TGT CGA AGC CAA TTG ACC AGC	208
N R Q H A R L R V C Q K I E K L *	748
AAT CGG CAG CAT GCT CGA TTA AGA GTA TGC CAA AAA ATA GAA AAG CTA TAA	225
ATATTTCAAAATAAAGAAGAATCCACATTGCAAAAAAAAAAAAAAAAAAAAA	799
	848

Figure 3A

CCGACGCGTGGGCGGCACGGTTTCGTGGGGACCCAGGCTTGCAAAGTGACGGTCATTTTCTCTTTCTTCTCCCTCTT 79
M M A L G A A G A T R V F V A M 16
GAGTCCTTCTGAG ATG ATG GCT CTG GGC GCA GCG GGA GCT ACC CGG GTC TTT GTC GCG ATG 140
V A A A L G G H P L L G V S A T L N S V 36
GTA GCG GCG GCT CTC GGC GGC CAC CCT CTG CTG GGA GTG AGC GCC ACC TTG AAC TCG GTT 200
L N S N A I K N L P P P L G G A A G H P 56
CTC AAT TCC AAC GCT ATC AAG AAC CTG CCC CCA CCG CTG GGC GGC GCT GCG GGG CAC CCA 260
G S A V S A A P G I L Y P G G N K Y Q T 76
GGC TCT GCA GTC AGC GCC GCG CCG GGA ATC CTG TAC CCG GGC GGC AAT AAG TAC CAG ACC 320
I D N Y Q P Y P C A E D E E C G T D E Y 96
ATT GAC AAC TAC CAG CCG TAC CCG TGC GCA GAG GAC GAG GAG TGC GGC ACT GAT GAG TAC 380
C A S P T R G G D A G V Q I C L A C R K 116
TGC GCT AGT CCC ACC CGC GGA GGC GAC GCA GGC GTG CAA ATC TGT CTC GCC TGC AGG AAG 440
R R K R C M R H A M C C P G N Y C K N G 136
CGC CGA AAA CGC TGC ATG CGT CAC GCT ATG TGC TGC CCC GGC AAT TAC TGC AAA AAT GGA 500
I C V S S D Q N H F R G E I E E T I T E 156
ATA TGC GTG TCT TCT GAT CAA AAT CAT TTC CGA GGA GAA ATT GAG GAA ACC ATC ACT GAA 560
S F G N D H S T L D G Y S R R T T L S S 176
AGC TTT GGT AAT GAT CAT AGC ACC TTG GAT GGC TAT TCC AGA AGA ACC ACC TTG TCT TCA 620
K M Y H T K G Q E G S V C L R S S D C A 196
AAA ATG TAT CAC ACC AAA GGA CAA GAA GGT TCT GTT TGT CTC CGG TCA TCA GAC TGT GCC 680
S G L C C A R H F W S K I C K P V L K E 216
TCA GGA TTG TGT TGT GCT AGA CAC TTC TGG TCC AAG ATC TGT AAA CCT GTC CTG AAA GAA 740
G Q V C T K H R R K G S H G L E I F Q R 236
GGT CAA GTG TGT ACC AAG CAT AGG AGA AAA GGC TCT CAT GGA CTA GAA ATA TTC CAG CGT 800
C Y C G E G L S C R I Q K D H H Q A S N 256
TGT TAC TGT GGA GAA GGT CTG TCT TGC CGG ATA CAG AAA GAT CAC CAT CAA GCC AGT AAT 860
S S R L H T C Q R H * 267
TCT TCT AGG CTT CAC ACT TGT CAG AGA CAC TAA 893
ACCAGCTATCCAAAATGCAGTGAACCTCTTTTATATAATAGATGCTATGAAAACCTTTTATGACCTTCATCAACTCAAT 972
CCTAAGGATATACAAGTTCTGTGGTTTCAGTTAAGCATTCCTCAATAACACCTTCCAAAAACCTGGAGTCTAAGAGCTTTG 1051
TTTCTTTATGGAACCTCCCCTGTGATTGCAGTAAATTACTGTATTGTAAATTCTCAGTGTGGCACTTACCTGTAAATGCA 1130
ATGAAACTTTTAATTATTTTCTAAAGGTGCTGCAC GGCCTATTTTCTCTGTATGTAAATTTTGTACACATTGA 1209
TTGTTATCTTGACTGACAAATATTCTATATTGAACTGAAGTAAATCATTTTCAGCTTATAGTTCTTAAAGCATAACCCCT 1288
TTACCCCATTTNATTCTAGAGTCNAGAACGCAAGGATCTCTTGAATGACAAATGATAGGTACCTAAATGTAAACATGA 1367

Figure 3B

AAATACTAGCTTATTTTCTGAAATGTACTATCTTAATGCTTAAATTATATTTCCCTTTAGGCTGTGATAGTTTTTGAAA 1446
TAAAATTTAACATTTAATATCATGAAATGKTATAAGTAGACATAAAAAAAAAAAAAAAAAAAGGGCGGCCGCTAGA 1525
CTAG 1529

Figure 4

E	F	G	T	R	V	G	R	Y	C	H	S	P	H	Q	G	S	S	A	C		20
GAA	TTC	GGC	ACG	AGG	GTT	GGG	AGG	TAT	TGC	CAC	AGT	CCC	CAC	CAA	GGA	TCA	TCG	GCC	TGC		60
M	V	C	R	R	K	K	K	R	C	H	R	D	G	M	C	C	P	S	T		40
ATG	GTG	TGT	CGG	AGA	AAA	AAG	AAG	CGC	TGC	CAC	CGA	GAT	GGC	ATG	TGC	TGC	CCC	AGT	ACC		120
R	C	N	N	G	I	C	I	P	V	T	E	S	I	L	T	P	H	I	P		60
CGC	TGC	AAT	AAT	GGC	ATC	TGT	ATC	CCA	GTT	ACT	GAA	AGC	ATC	TTA	ACC	CCT	CAC	ATC	CCG		180
A	L	D	G	T	R	H	R	D	R	N	H	G	H	Y	S	N	H	D	L		80
GCT	CTG	GAT	GGT	ACT	CGG	CAC	AGA	GAT	CGA	AAC	CAC	GGT	CAT	TAC	TCA	AAC	CAT	GAC	TTG		240
G	W	Q	N	L	G	R	P	H	T	K	M	S	H	I	K	G	H	E	G		100
GGA	TGG	CAG	AAT	CTA	GGA	AGA	CCA	CAC	ACT	AAG	ATG	TCA	CAT	ATA	AAA	GGG	CAT	GAA	GGA		300
D	P	C	L	R	S	S	D	C	I	E	G	F	C	C	A	R	H	F	W		120
GAC	CCC	TGC	CTA	CGA	TCA	TCA	GAC	TGC	ATT	GAA	GGG	TTT	TGC	TGT	GCT	CGT	CAT	TTC	TGG		360
T	K	I	C	K	P	V	L	H	Q	G	E	V	C	T	K	Q	R	K	K		140
ACC	AAA	ATC	TGC	AAA	CCA	GTG	CTC	CAT	CAG	GGG	GAA	GTC	TGT	ACC	AAA	CAA	CGC	AAG	AAG		420
G	S	H	G	L	E	I	F	Q	R	C	D	C	A	K	G	L	S	C	K		160
GGT	TCT	CAT	GGG	CTG	GAA	ATT	TTC	CAG	CGT	TGC	GAC	TGT	GCG	AAG	GGC	CTG	TCT	TGC	AAA		480
V	W	K	D	A	T	Y	S	S	K	A	R	L	H	V	C	Q	K	I	*		180
GTA	TGG	AAA	GAT	GCC	ACC	TAC	TCC	TCC	AAA	GCC	AGA	CTC	CAT	GTG	TGT	CAG	AAA	ATT	TGA		540
TCACCATTGAGGAACATCATCAATTGCAGACTGTGAAGTTGTGTATTTAATGCATTATAGCATGGTGAAAAATAAGGTT																					619
CAGATGCAGAAGAATGGCTAAAATAAGAAACGTGATAAGAATATAGATGATCACAAAAAAAAAAAAAAAAAAGATGCCG																					698
CCGC																					702

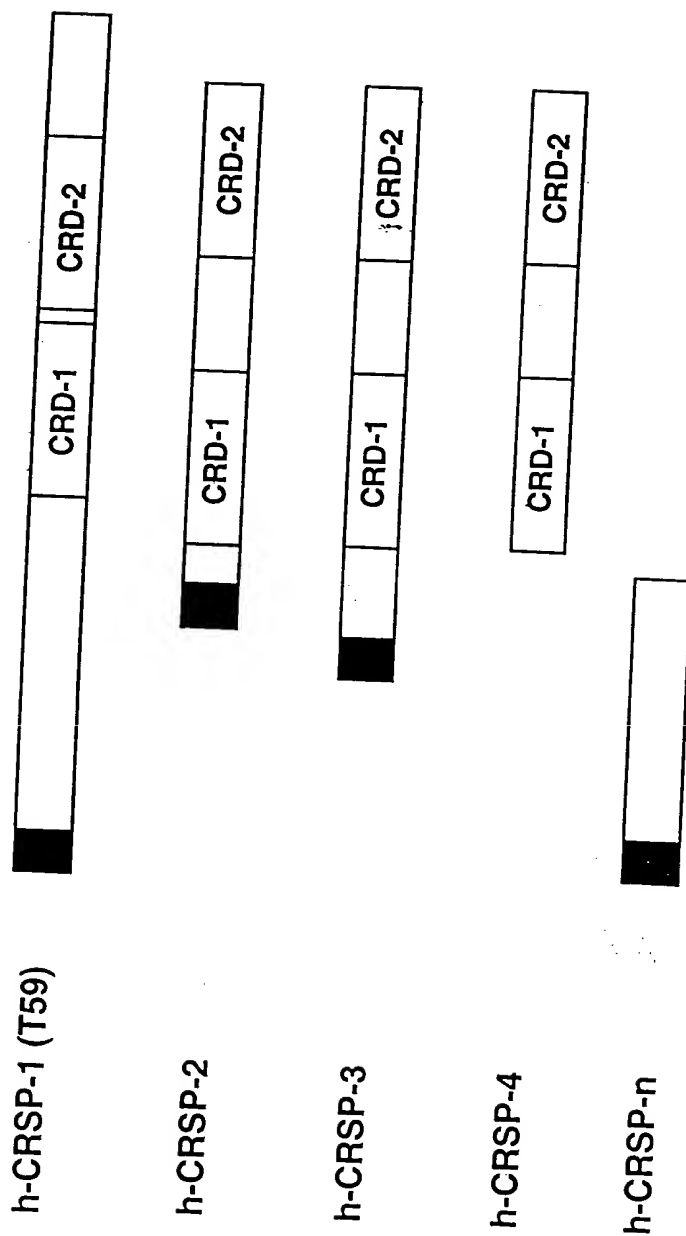
Figure 5

CTCGAGGCCAAAATTCGGCACGAGGCCGGGCTGTGGTCTAGCATAAAGGCGGAGCCCAGAAGAAGGGGCGGGGT	M	1
ATG		77
G E A S P P A P A R R H L L V L L L L L		21
GGA GAA GCC TCC CCA CCT GCC CCC GCA AGG CGG CAT CTG CTG GTC CTG CTG CTG CTC CTC		137
S T L V I P S A A A P I H D A D A Q E S		41
TCT ACC CTG GTG ATC CCC TCC GCT GCA GCT CCT ATC CAT GAT GCT GAC GCC CAA GAG AGC		197
S L G L T G L Q S L L Q G F S R L F L K		61
TCC TTG GGT CTC ACA GGC CTC CAG AGC CTA CTC CAA GGC TTC AGC CGA CTT TTC CTG AAA		257
G N L L R G I D S L F S A P M D F R G L		81
GGT AAC CTG CTT CGG GGC ATA GAC AGC TTA TTC TCT GCC CCC ATG GAC TTC CGG GGC CTC		317
P G N Y H K E E N Q E H Q L G N N T L S		101
CCT GGG AAC TAC CAC AAA GAG GAG AAC CAG GAG CAC CAG CTG GGG AAC AAC ACC CTC TCC		377
S H L Q I D K M T D N K T G E V L I S E		121
AGC CAC CTC CAG ATC GAC AAG ATG ACC GAC AAC AAG ACA GGA GAG GTG CTG ATC TCC GAG		437
N V V A S I Q P A E G S F E G D L K V P		141
AAT GTG GTG GCA TCC ATT CAA CCA GCG GAG GGG AGC TTC GAG GGT GAT TTG AAG GTA CCC		497
R M E E K E A L V P I Q K A T D S F H T		161
AGG ATG GAG GAG AAG GAG GCC CTG GTA CCC ATC CAG AAG GCC ACG GAC AGC TTC CAC ACA		557
E L H P R V A F W I I K L P R R R S H Q		181
GAA CTC CAT CCC CGG GTG GCC TTC TGG ATC ATT AAG CTG CCA CGG CGG AGG TCC CAC CAG		617
D A L E G G H W L S E K R H R L Q A I R		201
GAT GCC CTG GAG GGC GGC CAC TGG CTC AGC GAG AAG CGA CAC CGC CTG CAG GCC ATC CGG		677
D G L R K G T H K D V L E E G T E S S S		221
GAT GGA CTC CGC AAG GGG ACC CAC AAG GAC GTC CTA GAA GAG GGG ACC GAG AGC TCC TCC		737
H S R L S P R K T H L L Y I L R P S R Q		241
CAC TCC AGG CTG TCC CCC CGA AAG ACC CAC TTA CTG TAC ATC CTC AGG CCC TCT CGG CAG		797
L *		243
CTG TAG		803
GGGTGGGGACCGGGGAGCACCTGCCTGTAGCCCCCATCAGACCCTGCCCAAGCACCATATGGAAATAAAGTTCTTTCT		882
TACATCTAAAAAAAAAAAAAAAAAAAAAAAAAATTTGGCGGCCGC		928

406

Figure 7

The Human CRSP Family



FGT CGA C C C A C G C G T C O G C T G T G G C A G C C C A G C T A C C G G T C G T G A C C A G A T C C A G C T T G C A G C T C A G C T T T G T T C A T T C																										79
M Q R L G G I L L C T L																										12
GAATTGGGCGGCGGCCAGCGCGGAACAAAC ATG CAG CGG CTC GGG GGT ATT TTG CTG TGT ACA CTG																										145
L A A A V P T A P A P S P T V T W T P A	32																									
CTG GCG GCG GCG GTC CCC ACT GCT CCT GCT CCT TCC CCG ACG GTC ACT TGG ACT CCG GCG	205																									
E P G P A L N Y P Q E E A T L N E M F R	52																									
GAG CCG GGC CCA GCT CTC AAC TAC CCT CAG GAG GAA GCT ACG CTC AAT GAG ATG TTT CGA	265																									
E V E E L M E D T Q H K L R S A V E E M	72																									
GAG GTG GAG GAG CTG ATG GAA GAC ACT CAG CAC AAA CTG CGC AGT GCC GTG GAG GAG ATG	325																									
E A E E A A A K T S S E V N L A S L P P	92																									
GAG GCG GAA GAA GCA GCT GCT AAA ACG TCC TCT GAG GTG AAC CTG GCA AGC TTA CCT CCC	385																									
N Y H N E T S T E T R V G N N T V H V H	112																									
AAC TAT CAC AAT GAG ACC AGC ACG GAG ACC AGG GTG GGA AAT AAC ACA GTC CAT GTG CAC	445																									
Q E V H K I T N N Q S G Q V V F S E T V	132																									
CAG GAA GTT CAC AAG ATA ACC AAC AAC CAG AGT GGA CAG GTG GTC TTT TCT GAG ACA GTC	505																									
I T S V G D E E G K R S H E C I I D E D	152																									
ATT ACA TCT GTA GGG GAT GAA GAA GGC AAG AGG AGC CAT GAA TGT ATC ATT GAT GAA GAC	565																									
C G P T R Y C Q F S S F K Y T C Q P C R	172																									
TGT GGG CCC ACC AGG TAC TGC CAG TTC TCC AGC TTC AAG TAC ACC TGC CAG CCA TGC CGG	625																									
D Q Q M L C T R D S E C C G D Q L C A W	192																									
GAC CAG CAG ATG CTA TGC ACC CGA GAC AGT GAG TGC TGT GGA GAC CAG CTG TGT GCC TGG	685																									
G H C T Q K A T K G G N G T I C D N Q R	212																									
GGT CAC TGC ACC CAA AAG GCC ACC AAA GGT GGC AAT GGG ACC ATC TGT GAC AAC CAG AGG	745																									
D C Q P G L C C A F Q R G L L F P V C T	232																									
GAT TGC CAG CCT GGC CTG TGT TGT GCC TTC CAA AGA GGC CTG CTG TTC CCC GTG TGC ACA	805																									
P L P V E G E L C H D P T S Q L L D L I	252																									
CCC CTG CCC GTG GAG GGA GAG CTC TGC CAT GAC CCC ACC AGC CAG CTG CTG GAT CTC ATC	865																									
T W E L E P E G A L D R C P C A S G L L	272																									
ACC TGG GAA CTG GAG CCT GAA GGA GCT TTG GAC CGA TGC CCC TGC GCC AGT GGC CTC CTA	925																									
C Q P H S H S L V Y M C K P A F V G S H	292																									
TGC CAG CCA CAC AGC CAC AGT CTG GTG TAC ATG TGC AAG CCA GCC TTC GTG GGC AGC CAT	985																									
D H S E E S Q L P R E A P D E Y E D V G	312																									
GAC CAC AGT GAG GAG AGC CAG CTG CCC AGG GAG GCC CCG GAT GAG TAC GAA GAT GTT GGC	1045																									
F I G E V R Q E L E D L E R S L A Q E M	332																									
TTC ATA GGG GAA GTG CGC CAG GAG CTG GAA GAC CTG GAG CGG AGC CTA GCC CAG GAG ATG	1105																									

Figure 8B

A F E G P A P V E S L G G E E E I *
GCA TTT GAG GGG CCT GCC CCT GTG GAG TCA CTA GGC GGA GAG GAG GAG ATT TAG 350
1159
GCCCAGACCCAGCTGAGTCACTGGTAGATGTGCAATAGAAATGGCTAATTTATTTTCCCAGGAGTGTCCCCAAGTGTGG 1238
AATGGCCGCAGCTCCTTCCCAGTAGCTTTTCTCTGGCTTGACAAGGTACAGTGCAGTACATTTCTTCCAGCCGCCCTG 1317
CTTCTCTGACTTGGGAAAGACAGGCATGGCGGGTAAGGGCAGCGGTGAGTGCCTCCCTCGCTGTTGCTAGAAACGCTGTC 1396
TGTTCTTTCATGGATGGAAGATTTGTTTGAAGGGAGAGGATGGGAAGGGGTGAAGTCTGCTCATGATGGATTGGGGGA 1475
TACAGGGAGGAGGATGCCTGCCTTGCAGACGTGGACTTGGCAAAATGTAACCTTTGCTTTTGTCTTGCGCCGCTCCCAT 1554
GGGCTGAGGCAGTGGCTACACAAGAGCTATGCTGCTCTGTGGCCTCCACATATTTCATCCCTGTGTTTCAGCTCCTACC 1633
TCACTGTGAGCACAGCCCTTCATAGCCACGCCCCCTCTTGCTCACCACAGCCTAGGAGGGGACCAGAGGGGACTTCTCT 1712
CAGAGCCCCATGCTCTCTCTCTCAACCCCATACCAGCCTCTGTGCCAGCGACAGTCCTTCCAAATGGAGGGAGTGAAAT 1791
CCTTTGGTTTAATTATTTTCTCCTTCAAGGCACGCCTGCCACTAAGGTCAGGCTGACTTGTCATGTCCCTCTAACGTTCCG 1870
TAGCAGTGTGGTGGACACTGTCTTCCACCGACTGCTTCAATACCTCTGAAAGCCAGTGTCTCGGAGTGCAGTTCGTGTAA 1949
ATTAATTTGCAGGAAGTATACTTGGCTAATTGTAGGGCTAGGATTGTGAATGAAATTTGCAAAGTCGCTTAGCAACAAT 2028
GGAAAGCCTTTCTCAGTCACACCGAGAAGTCACAACCAAGCCAGGTTGTGTAGAGTACAGCTGTGACATACAGACAGAA 2107
GAAGGCTGGGCTGGATGTCAGGCCTCAGATGACGGTTTCAGGTGCCAGGAACCTATTACCATTCTGTATCTATCCAGAGT 2186
TATTAAAATTGAAAGTTGCACACATTTGTATAAGCATGCCTTTCTCCTGAGTTTAAATTATATGTATACACAAACATG 2265
TGGCCCTCAAAGATCATGCACAAACCACTACTCTTTGCTAATTCTTGGACTTTTCTCTTTGATTTTCAATAAATACAAA 2344
TCCCCTTCATGCAAAAAAAAAAAAAAAGGGCGGCCGC 2381